

GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:09:48 ; Search time 1470 Seconds
(without alignments)
13572.847 Million cell

updates/sec

Title: US-09-300-482-225
Perfect score: 316
Sequence: 1
gataaggtgcgcacactgag.....gaatggaggaggagtacttcga 316

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 111716937 seqs, 31560637849 residues

Total number of hits satisfying chosen parameters: 223433874

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_NA_Main:*
1: /ABSS/Data/CRF/ptodata/1/pna/PCTUSA_COMB.seq:*
2: /ABSS/Data/CRF/ptodata/1/pna/PCTUSB_COMB.seq:*
3: /ABSS/Data/CRF/ptodata/1/pna/PCTUSC_COMB.seq:*
4: /ABSS/Data/CRF/ptodata/1/pna/PCTUSD_COMB.seq:*
5: /ABSS/Data/CRF/ptodata/1/pna/US075_COMB.seq:*
6: /ABSS/Data/CRF/ptodata/1/pna/US076_COMB.seq:*
7: /ABSS/Data/CRF/ptodata/1/pna/US077_COMB.seq:*
8: /ABSS/Data/CRF/ptodata/1/pna/US078_COMB.seq:*
9: /ABSS/Data/CRF/ptodata/1/pna/US079_COMB.seq:*
10: /ABSS/Data/CRF/ptodata/1/pna/US080_COMB.seq:*
11: /ABSS/Data/CRF/ptodata/1/pna/US081_COMB.seq:*
12: /ABSS/Data/CRF/ptodata/1/pna/US082_COMB.seq:*
13: /ABSS/Data/CRF/ptodata/1/pna/US083_COMB.seq:*
14: /ABSS/Data/CRF/ptodata/1/pna/US084_COMB.seq:*
15: /ABSS/Data/CRF/ptodata/1/pna/US085_COMB.seq:*